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MPsrch\_pp protein - protein database search, using Smith-Waterman algorithm

Tabular output Run on: not generated Sat May 13 10:16:40 2000; MasPar time 4.56 Seconds 327.460 Million cell updates/sec

Description: Perfect Score: Title:

Sequence: >US-09-331-631-3 \(186-248\) from US09331631.pep (5 of 5) 492 1 KRDPQOREYEDCRRRCEQQE.....LINPQRGGSGRYEEGEEKQS

Scoring table: PAM 150 Gap 11

Searched:

Post-processing:

188963 seqs,

23686106 residues

Minimum Match Listing first 4 0 8 5 summaries

Database: a-geneseq35 1:geneseqp

Statistics: Mean 25.162; Variance 101.062; scale 0.249

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

## SUMMARIES

2222248454432333333333333333333333333333	Result No.
1111446 1117488 1117788 1117788 1117788 1117788 1117788 1117788 111778 117778 1177	Score
93.9 91.3 36.2 37.4 38.2 20.9 117.1 117.1 117.1 116.9 116.9 116.9 116.9 116.9	Query Match
222566625 5666655 221119993 566665 266893 266893 266893 266893	1
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W62838 W62838 W62838 W62838 R20181 R20183 R12223 W12223 R142823 R14283 R14283 R14283 R14283 R14283 R1438 R14383 R14383 R14383 R14383 R14383 R14383 R14383 R14383 R14383 R14383 R14383 R14383 R14383 R14383 R14383 R14383 R14383 R14383 R1	ID
Macadamia integrifolia Macadamia integrifolia Theobroma cacao antimi Sequence encoded by 67 Gossypium hirsutum ant Hordeum vulgare antimi Zea mays antimicrobial Human androgen recepto Mycobacterium species Mycobacterium species Protein derived from 7 Zea mays antimicrobial maize pe Human androgen recepto Stenocarpus sinuatus a Spinocerebellar ataxia Soybean glycinin A3B4 Mycobacterium tubercul Human secreted protein Human secreted protein	Description
8.17e-37 1.79e-35 2.98e-08 2.98e-08 3.23e-04 2.37e-01 2.08e+00 2.08e+00 2.08e+00 7.99e+00 7.99e+00 9.66e+00 9.66e+00 9.66e+01 1.17e+01 1.17e+01 1.41e+01 1.41e+01 1.41e+01 1.70e+01 1.70e+01 1.70e+01 1.70e+01	Pred. No.

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Yeast transcriptional	Murine vascular endoth	Murine vascular endoth	Human vascular endothe	Heart vascular endothe	Human VEGF-B full leng	Human VEGF-B truncated	Mycobacterium species	A Drosophila apoptosis	Murine vascular endoth	Vascular endothelial g	Rat YT521 gene product	FMR-1 gene product.	Moloney murine leukemi	GST-HD fusion protein	GST-HD fusion protein	Osteoinductive retrovi					
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Best Local Similarity 100.0%;
Matches 63; Conservative
W62830 standard; Protein; 625 AA. W62830; W62830; 27-OCT-1998 (first entry) Macadamia integrifolia antimicrobial protein. antimicrobial protein; infestation; control.
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The sequence is that of an antimicrobial protein which can be used to control microbial infestations in plants and mammalian
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02-JUL-1998:
22-DEC-1997; AU0874.
4
20-DEC-1996; AU-004275.
(RETR-) COOP RES CENT TROPICAL PLANT PATHOLOGY.
BOWEY NI, GOULTER KC, Green JL, Manners JM, Mar
WPI; 98-377279/32.
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29. .666
/note= "mature protein"
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Pred. No. 6.47e-40;
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22-DEC-1997; AU0874.
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(RETR-) COOP RES CENT TR
Bower NI, Goulter KC, Gr
WPI; 98-377279/32.
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(RETR-) COOP RES CENT TROPIC

POWER NI GOULTER KC, Green

WPI: 98-377279/32.

N-PSDB: V42310.
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22-DEC-1997;
20-DEC-1996;
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The sequence is that of an antimicrobial protein which can be used to control microbial infestations in plants and mar
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The sequence is that of an antimicrobial protein which can be used to control microbial infestations in plants and mammalian
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Macadamia integrifolia antimicrobial protein.
antimicrobial protein; infestation; control.
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EQS 248
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Similarity 93.7%;
59; Conservet'
                                                                                                                                                                Similarity
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ce= "mature protein"
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Green JL, Manners JM, Mar
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Green JL, Manners JM, Marcus
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Pred. No. 1.79e-35;
2; Mismatches 3
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Pred. No. 8.17e-37;
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Matches
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Claim 4; Fig 2; 59pp; English.
Claim 4; Fig 2; 59pp; English.
The inventors claim a 67 kD and 31 kD T. cacao protein, and fragments, and encoding DNAs. The 47 kD and 31 kD proteins are derived from the 67 kD precursor. T. cacao protein cDNA was detected in a cDNA library prepared from inmature cocoa beans RNA using a probe based on the AA sequence of a CNBr peptide common to the 47 kD and 31 kD polypeptides. Homology searches revealed clos homologies between the 67 kD polypeptide and the vicilins, which a
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11-JUN-1990; GB-013016.
(MRSC ) MARS UK LTD.
Spencer ME, Hodge R, Deakin
WPI; 92-024418/03.
                                                                                                                                                                         seed storage protes
requence 566 AA;
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Sequence encoded by 67 kD T
Cocoa; flavour; vicilin; see
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The sequence is that of an antimicrobial protein which can
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27-OCT-1998
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Local Similarity
nes 23; Conse
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   82 QRQYQQCQGRCQEQQQGQREQQQCQRKCWEQYKEQERGEHENYHNHKKNRSEEEEGQQRN 141
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                                                                           36.2%;
ilarity 37.7%;
Conservative
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larity 37.7%;
Conservative
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                                                                        Score 178;
Pred. No. 2.
20; Mismatc
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Pred. No. 2.98e-08;
20; Mismatches 15
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98e-08;
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JLT 8
W62835 standard;
W62835;
27-OCT-1998 (fir
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Novel anti-microbial protein from e.g. Macadamna uncey......

Novel anti-microbial protein infestations of plants or mammals useful for controlling microbial infestations of plants or mammals claim 1; Page 49-51; 96pp; English.

The sequence is that of an antimicrobial protein which can fine sequence is that of an antimicrobial protein which can antimicrobial infestations in plants and mammalian
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22-DEC-1997; AU0874.
20-DEC-1996; AU-004275.
20-DEC-1996; AU-004275.

(RETR-) COOP RES CENT TROPICAL PLANT PATHOLOGY.
BOWER NI, Goulter KC, Green JL, Manners JM, Mar
WPI; 98-377279/32.
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22-DEC-1997; AU0874.
20-DEC-1996; AU-004275.
(RETR-) COOP RES CENT TROPICAL PLANT PATHOLOGY.
BOWER NI, Goulter KC, Green JL, Manners JM, Mar
WPI; 98-377279/32.
                                                                                                                                                                                                                Novel anti-microbial protein from e.g. Macadamia integrifolia useful for controlling microbial infestations of plants or man Claim 1; Page 60-62; 96pp; English.

The sequence is that of an antimicrobial protein which can
                                                                                                                                                                                                                                                                                                                                                 Hordeum vulgare W09827805-A1.
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W62832 standard; Protein;
                                                                                                                                                                               Sequence
                                                                                                                                                                                            be used to control microbial infestations animals.
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nes 23; Conser
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                        Protein; 593
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Pred. No. 3.
13; Mismatc
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Pred. No. 5.58e-06;
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                                                                                                                           e 135; DB 1; L. No. 3.23e-04; Mismatches 16;
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Best Local
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30-MAY-1991.
19-OCT-1990; U06015.
17-NOV-1989; US-438775.
(ARCH-) ARCH DEV CORP.
Liao S, Chang C;
WPI; 91-178048/24.
                                                                                                                                                                                                                                                                                                                                                                              N-PSDB; Q12001.

Androgen receptor and TR2 DNA binding proteins - DNA sequences and antibodies for detection and quantification methods and antibodies for detection and quantification methods Claim 25; Fig 3; 79pp; English.

This sequence was deduced from a cDNA clone isolated by screening commercially available human testis and prostate lambda gtl1 cDNA libraries. The sequence is very similar to that of rat Ra and in the DNA-binding domain it is identical to that of rat pNA-binding domain. Homology comparisons with other known steroid receptors indicate that hAR is more closely related to glucocorticoid, mineralo-corticoid and progesterone receptors than to v-erb-A or treceptors for cestrogen, vitamin D and thyroid hormones.
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22-DEC-1997; /
20-DEC-1996; /
                                                 W14783;
W14783;
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(RETR-) COOP RES CENT TROPIC
BOWER NI, Goulter KC, Green
WPI; 98-377279/32.
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The sequence is that of an antimicrobial protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Zea mays antimicrobial antimicrobial protein; Zea mays. W09827805-A1.
                                                                                                                                                                                                                                                                                                                                                                         Sequence
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                         22-JUN-1997
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                                                                                                                                                                       202 EQQEPRQQYQCQRRCREQQRQHGRGGDLINPQRGGSGRYEEGEEKQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         196
                                                                                                                                                                                                      64 QQQQQQQQQQQETSPRQQQQQQGEDGSPQAHRRGPTG-YLVLDEEQ 108
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DNA-binding protein; steroid hormone
                                                                          standard; Protein;
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                                                                                                                                                                                                                                                                                                 Similarity
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                       (first entry)
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larity 32.7%;
Conservative
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/label= DNA-binding do
/note= "cysteine-rich"
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                                                                                                                                                                                                                                                                                                 18.7%;
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Pred. No. 2.:
17; Mismatcl
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                                                                                                                                                                                                                                                                         Score 92; DB 1; L
Pred. No. 2.08e+00;
10; Mismatches 18
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.37e-01
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Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                      DNA encoding androgen receptor protein - useful for transforming eukaryotic hosts for protein expression and subsequent antibody prodn. Disclosure; Fig. 4: 41pp; English.

Androgen receptor protein (AR) is used to produce mono- or poly-clonal antibodies. These are used for the detection and quantification of AR in the presence of endogenous androgen, as androgen will not interfere with binding. They may be used in assays to determine and quantify cellular distribution of AR in tumour tissue, and are esp. useful for evaluating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FGF genes - used to inhibit gene expression. For the treatment of benign prostatic hyperplasia Disclosure; Page 22-28; 51pp; English.

Human androgen receptor (W14783) binds testosterone and, acting at the transcriptional level, regulates the growth of normal prostatic cells. Antisense oligonucleotides (see also T63200, T63404-05) based on an androgen receptor CDNA clone (see also T63407) can be used to prevent androgen receptor gene expression, thereby inhibiting the growth or survival of prostatic cells for the treatment of benign prostatic hyperplasia and prostate cancer.
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W09711170-A1.
27-MAR-1997.
Y04866;
06-JUL-1999
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Y04866 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                   prostate cancers to determine responsiveness to androgen
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14-APR-1988; US-182646.
(UYNC-) University of North Carolina.
French FS, Wilson EM, Joseph DR, Lubahn
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20-SEP-1995; US-004018.
(WORC-) WORCESTER FOUND BIOMEDICAL RES
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llarity 37.0%;
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Claim 32; Fig 19C; 309pp; French.
Sequences Y04742-Y05000 and Y07201-Y07204 represent secreted proteins from various Mycobacterium species microorganisms. The encoding nucleotide sequences can be used as primers and probes for methods for detecting and identifying mycobacteria, especially belonging to the M. tuberculosis complex. The encoded proteins can be used in vaccines for immunisation against a bacterial or viral infection.
Sequence 112 AA;
                                                                                                                                                                                            N-PSDB; x34113.

N-PSDB; x34113.

Nycobacterial DNA vectors containing reporter constructs - for identifying coding or promoter sequences involved in infection-associated protein expression

Claim 32; Fig 19A; 309pp; French.

Sequences y04742-y05000 and y07201-y07204 represent secreted proteins from various Mycobacterium species microorganisms. The encoding nucleotide sequences can be used as primers and probes for methods
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Gicquel B, Li
                                                                                                                       for detecting and identifying mycobacteria, especially belonging the M. tuberculosis complex. The encoded proteins can be used invaccines for immunisation against a bacterial or viral infection
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Secreted protein; Mycobacterium; primer; PCR; amplification; hybridisation; detection; vaccine; immunisation; infection.
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WPI; 99-181
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local S
Matches 1
                                                                                                                                                                                                                                                  Query Match 17.1%;
Best Local Similarity 44.0%;
Matches 11; Conservative
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30-SEP-1996; E04254.
25-MAR-1996; DE-011758.
04-OCT-1995; DE-036890.
(BADI ) BASF AG.
(BADI ) BASF AG.
Kahmann R. Quadbeck-seeger C;
WPI: 97-226165/20.
N-PSDB; T62940.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      W14987;
W14987;
W14987;
W14987;
W14987;
W14987;
W2-FEB-1998 (first entry)
W2-FEB-1998 (first entry)
W3-FEB-1998 (first entry)
W4-11490 maydis.
W4-11490 maydis.
W4-11490 maydis.
                                                                                                                                                                                                                                                                                                                                                                                                                           02-JUL-1998.
02-DEC-1997; AU0874.
22-DEC-1996; AU-004275.
(RETR-) COOP RES CENT TROPICAL PLANT PATHOLOGY.
Bower NI, Goulter KC, Green JL, Manners JM, Marcus JP;
WPI; 98-377279/32.
Novel anti-microbial protein from e.g. Macadamia integrifolia -
useful for controlling microbial infestations of plants or mammals
Disclosure; Page 60; 96pp; English.
The sequence is that of an antimicrobial protein which can
be used to control microbial infestations in plants and mammalian
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2027 RDYERERERDGDLHDARDGRDGRYGDS 2053
|: :|:: | ||| ::: | ||| ::
217 REQQRQHGRGGGDLINPQRGGSGRYEEG 243
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New regulatory gene from Ustilago maydis and derived protein useful as target for identifying fungicides, also mutant fungus for infecting maize claim 9; Page 9-18; 42pp; German.

The gene is a regulatory gene in U. maydis (a maize pathogen) and probably has a repressive action.

Utilago strains with a mutation is this gene can form galls, but not spores, in maize, so the galls (used as food) are visibly detectable.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    27-OCT-1998 (first entry)
Zea mays antimicrobial protein.
antimicrobial protein; infestation; control.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /T 15
W62836 standard; Protein; 33 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                JT 14
W14987 standard; Protein; 2289 AA.
                                                                                                                                                                                                                                                                                                                                                                                Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                  animals
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Zea mays.
WO9827805-A1.
                                                                                                                       196 DCRRRC-EQQE--PRQYQCQRRCR 217
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               y Match 17.3%;
Local Similarity 37.0%;
hes 10; Conservative
                                                                                                                                                                                   6 ECRRQCLRRHEGQPWETQECMRRCR 30
                                                                                                                                                                                                                                                                                                                                                                                    33 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 85; DB 1; Length 2289; Pred. No. 7.99e+00; 9; Mismatches 8; Indels
                                                                                                                                                                                                                                           Score 84; DB 1; Length 33; Pred. No. 9.66e+00; 7; Mismatches 4; Indels
                                                                                                                                                                                                                                                     Indels
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